Search term = sucrose transporter

Sucrose transpoter coding sequences available in EMBL or SwissProt databases prior to August 07, 1997



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Description:transporter*]) | [libs={embl swissprot}-Description:**sucrose** transporter*]) & [libs-DateCreated#19000101:19970806]) " found 5 entries Query "((([libs={embl swissprot}-Description:sucrose*] & [libs={embl swissprot}-

EMBL	T EMBL:LESUTI	T EMBL:N
-	ESUTI	TSUTIA
D	ESUTI	NTSUTIA
AccNumber Molecule Description	X82275 R	X82276 R
olecule	X	A
Description	H :	EMBL:NTSUTIA NTSUTIA X82276 RNA N.tabacum SUT1a mRNA for sucrose transporter
Organism	Lycopersicon esculentum (tomato) Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Asteridae; lamiids; Solanales; Solanaceae; Solanum; Lycopersicon.	Nicotiana tabacum (common tobacco) Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta;
DateCreated	28-OCT-1994	28-OCT-1994

Search term = sucrose permease

Sucrose transpoter coding sequences available in EMBL or SwissProt databases prior to August 07, 1997



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Description:permease*]) | [libs={embl swissprot}-Description:sucrose permease*]) & [libs-DateCreated#19000101:19970807]) " found 12 entries Query "((([libs={embl swissprot}-Description:sucrose*] & [libs={embl swissprot}-

Swiss-Prot:PTSA_LACLA	Swiss-Prot:CSCB_ECOL1	Swiss-Prot EMBL
PTSA_LACLA Q04938	CSCB_ECOLI P30000	I
<u>Q04938</u>	P30000	Acc Number
		Molecule
PTS system, sucrose- specific IIABC component (EIIABC- SCR) (Sucrose- permease IIABC component)	Sucrose transport protein (Sucrose permease).	Description
LACTOCOCCUS LACTIS (SUBSP. LACTIS) (STREPTOCOCCUS LACTIS). BACTERIA; FIRMICUTES; LACTOBACILLALES; STREPTOCOCCACEAE; LACTOCOCCUS.	ESCHERICHIA COLI. BACTERIA; BACTERIA; PROTEOBACTERIA; GAMMAPROTEOBACTERIA; ENTEROBACTERIALES; ENTEROBACTERIACEAE; ESCHERICHIA.	Organism
01-JUN-1994	01-APR-1993	DateCreated

Swiss-Prot:PTSB_BACSU P	Swiss-Prot.PTSA_STRMU P	Swiss-Prot.PTSA_PEDPE P	
PTSB_BACSU	PTSA_STRMU <u>P12655</u>	PTSA_PEDPE	
P05306	<u>P12655</u>	<u>P43470</u>	
PTS system, sucrose- specific IIBC component (EIIBC-SCR) (Sucrose- permease IIBC component) (Phosphotransferase	PTS system, sucrose-specific IIABC component (EIIABC-SCR) (Sucrose-permease IIABC component) (Phosphotransferase enzyme II, ABC component) (EC 2.7.1.69) (EII-SCR).	PTS system, sucrose-specific IIABC component (EIIABC-SCR) (Sucrose-permease IIABC component) (Phosphotransferase enzyme II, ABC component) (EC 2.7.1.69) (EII-SCR).	enzyme II, ABC component) (EC 2.7.1.69) (EII-SCR) (Fragment).
BACILLUS SUBTILIS. BACTERIA; FIRMICUTES; BACILLALES; BACILLACEAE; BACILLUS.	STREPTOCOCCUS MUTANS. 01-OCT-1989 BACTERIA; FIRMICUTES; LACTOBACILLALES; STREPTOCOCCACEAE; STREPTOCOCCUS.	PEDIOCOCCUS PENTOSACEUS. BACTERIA; FIRMICUTES; LACTOBACILLALES; LACTOBACILLACEAE; PEDIOCOCCUS.	
01-NOV-1988	01-OCT-1989	01-NOV-1995	

٦	7	7
Swiss-Prot:PTSB_STAXY	Swiss-Prot:PTSB_SALTY	Swiss-Prot:PTSB_KLEPN
PTSB_STAXY	PTSB_SALTY	PTSB_KLEPN
P51184	P08470	P27219
PTS system, sucrose- specific IIBC component (EIIBC-SCR) (Sucrose- permease IIBC component) (Phosphotransferase enzyme II, BC component) (EC 2.7.1.69) (EII-SCR).	PTS system, sucrose-specific IIBC component (EIIBC-SCR) (Sucrose-permease IIBC component) (Phosphotransferase enzyme II, BC component) (EC 2.7.1.69) (EII-SCR).	enzyme II, BC component) (EC 2.7.1.69) (EII-SCR). PTS system, sucrose-specific IIBC component (EIIBC-SCR) (Sucrose-permease IIBC component) (Phosphotransferase enzyme II, BC component) (EC 2.7.1.69) (EII-SCR).
STAPHYLOCOCCUS XYLOSUS. BACTERIA; FIRMICUTES; BACILLALES; STAPHYLOCOCCUS.	SALMONELLA TYPHIMURIUM. BACTERIA; PROTEOBACTERIA; GAMMAPROTEOBACTERIALES; ENTEROBACTERIACEAE; SALMONELLA.	KLEBSIELLA PNEUMONIAE. BACTERIA; PROTEOBACTERIA; GAMMAPROTEOBACTERIALES; ENTEROBACTERIACEAE; KLEBSIELLA.
01-OCT-1996	01-AUG-1988	01-AUG-1992

EMBL:SOS21		Swiss-Prot:STP_SPIOL	Swiss-Prot.PTSB_VIBAL
SOS21		STP_SPIOL	PTSB_VIBAL
X67125 S51273		<u>Q03411</u>	P22825
RNA			
S.oleraceae mRNA for sucrose permease		Sucrose transport protein (Sucrose permease) (Sucrose-proton symporter).	PTS system, sucrose- specific IIBC component (EIIBC-SCR) (Sucrose- permease IIBC component) (Phosphotransferase enzyme II, BC component) (EC 2.7.1.69) (EII-SCR).
Spinacia oleracea (spinach) Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Caryophyllidae; Caryophyllales;	TRACHEOPHYTA; SPERMATOPHYTA; SPERMATOPHYTA; MAGNOLIOPHYTA; EUDICOTYLEDONS; CORE EUDICOTS; CARYOPHYLLALES; AMARANTHACEAE; SPINACIA.	SPINACIA OLERACEA (SPINACH). (SPINACH). EUKARYOTA; VIRIDIPLANTAE; STREPTOPHYTA; FMBRYOPHYTA:	VIBRIO ALGINOLYTICUS. BACTERIA; PROTEOBACTERIA; GAMMAPROTEOBACTERIA; VIBRIONALES; VIBRIONACEAE; VIBRIO.
17-DEC-1992		01-JUN-1994	01-AUG-1991

EMBL:VFZ93774 VFZ93774 Z93774 RNA	EMBL:STSUCTR STSUCTR X69165 RNA	EMBL:NTSUTIB NTSUTIB X82277 RNA	***************************************
3774 Z937	CTR X691	TIB X822	
74 RNA	65 RNA	277 RNA	
V.faba mRNA for sucrose transporter	S.tuberosum mRNA for sucrose transporter	N.tabacum SUT1b mRNA for sucrose transporter	
Vicia faba (fava bean) Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids I; Fabales; Fabaceae; Papilionoideae; Vicicae; Vicia.	Solanum tuberosum (potato) Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Asteridae; lamiids; Solanales; Solanaceae; Solanum.	Nicotiana tabacum (common tobacco) Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Asteridae; lamiids; Solanales; Solanaceae; Nicotiana.	lamiids; Solanales; Solanaceae; Nicotiana.
09-APR-1997	23-DEC-1993	28-OCT-1994	

	Г <u>EMBL:ECCSCI</u>	The second secon
	.SCB	
	ECCSCB	Tanana anno 1
	X63740 DNA S47896	
	E.coli cscB gene for Sucrose permease	
Gammaproteobacteria; Enterobacteriales; Enterobacteriaceae; Escherichia.	3 3 3 3	Chenopodiaceae; Spinacia.
•	13-MAR-1992	

SRS 7.0.2 | feedback

Search term = sucrose phosphotransferase

Sucrose transpoter coding sequences available in EMBL or SwissProt databases prior to August 07, 1997



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phosphotransfera se*]) & [libs-DateCreated#19000101:19970807]) " found 12 entries Query "((([libs={embl swissprot}-Description:sucrose*] & [libs={embl swissprot}-Description:phosphotransferase*]) | [libs={embl swissprot}-Description:sucrose

	7	
EMBL:BSSACP	EMBL:TBAA55334	EMBI. Swiss-Prot
BSSACP	TBAA55334	IE
J03006 M33761	AA255334 RNA	AccNumber
DNA	RNA	Molecule
B.subtilis sacT gene, complete cds, enzyme II Bacteria; Firmicutes; [sacP) gene (of the sucrose phosphotransferase phosphotransferase system), complete cds,	T3538 Bloodstream form of serodeme ILTat1.1 Trypanosoma brucei brucei cDNA 5' similar to gp J03006 Sac genesucrose phosphotransferase system [B.subtilis], mRNA sequence.	Description
Bacillus subtilis Bacteria; Firmicutes; Bacillales; Bacillaceae; Bacillus.	Trypanosoma brucei brucei Eukaryota; Euglenozoa; Kinetoplastida; Trypanosomatidae; Trypanosoma; Trypanosoma; Trypanosoma	Organism
16-JUL-1988	18-MAR-1997	DateCreated

٦	7	7	7
Swiss-Prot:PTSA_PEDPE	Swiss-Prot:PTSA_LACLA	EMBL:S74697	EMBL:BSU34876
PTSA_PEDPE	PTSA_LACLA Q04938	S74697	BSU34876
P43470	<u>Q04938</u>	S74697	U34876
		RNA	DNA
PTS system, sucrose- specific IIABC component PENTOSACEUS (EIIABC-SCR) (Sucrose- permease IIABC component) (Phosphotransferase enzyme II, ABC	PTS system, sucrose- specific IIABC component (SUBSP. LACTIS) (EIIABC-SCR) (Sucrose- permease IIABC component) (Phosphotransferase enzyme II, ABC component) (EC 2.7.1.69) (EII-SCR) (Fragment).	4All antigen=putative sucrose-specific phosphotransferase enzyme II homolog [mice, testis, mRNA Partial, 775 nt].	and sucrase gene, 5' end. Bacillus stearothermophilus phosphotransferase system sucrose enzyme II (surP) gene, complete cds.
PEDIOCOCCUS PENTOSACEUS. BACTERIA; FIRMICUTES; LACTOBACILLALES; LACTOBACILLACEAE; PEDIOCOCCUS.	LACTOCOCCUS LACTIS (SUBSP. LACTIS) (STREPTOCOCCUS LACTIS). BACTERIA; FIRMICUTES; LACTOBACILLALES; STREPTOCOCCACEAE; LACTOCOCCUS.	Mus sp. Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.	Geobacillus stearothermophilus Bacteria; Firmicutes; Bacillales; Bacillaceae; Geobacillus.
01-NOV-1995	01-JUN-1994	13-MAY-1995	19-FEB-1997

Swiss-Prot:PTSB_KLEPN	Swiss-Prot:PTSB_BACSU	Swiss-Prot:PTSA_STRMU	
PTSB_KLEPN P27219	PTSB_BACSU P05306	PTSA_STRMU P12655	
PTS system, sucrose- specific IIBC component (EIIBC-SCR) (Sucrose- permease IIBC component) (Phosphotransferase enzyme II, BC component) (EC <u>2.7.1.69</u>) (FII-SCR).	PTS system, sucrose-specific IIBC component (EIIBC-SCR) (Sucrose-permease IIBC component) (Phosphotransferase enzyme II, BC component) (EC 2.7.1.69) (EII-SCR).	PTS system, sucrose- specific IIABC component (EIIABC-SCR) (Sucrose- permease IIABC component) (Phosphotransferase enzyme II, ABC component) (EC 2.7.1.69) (EII-SCR).	component) (EC <u>2.7.1.69</u>) (EII-SCR).
KLEBSIELLA PNEUMONIAE. BACTERIA; BROTEOBACTERIA; PROTEOBACTERIA; GAMMAPROTEOBACTERI A; ENTEROBACTERIALES; ENTEROBACTERIACEAE; KLEBSIELLA.	BACILLUS SUBTILIS. BACTERIA; FIRMICUTES; BACILLALES; BACILLACEAE; BACILLUS.	STREPTOCOCCUS MUTANS. BACTERIA; FIRMICUTES; LACTOBACILLALES; STREPTOCOCCACEAE; STREPTOCOCCUS.	
01-AUG-1992	01-NOV-1988	01-OCT-1989	

Swiss-Prot:PTSB_VIBAL	Swiss-Prot:PTSB_STAXY	Swiss-Prot:PTSB_SALTY
PTSB_VIBAL P22825	PTSB_STAXY <u>P51184</u>	PTSB_SALTY P08470
PTS system, sucrose- specific IIBC component (EIIBC-SCR) (Sucrose- permease IIBC component) (Phosphotransferase enzyme II, BC component) (EC 2.7.1.69) (EII-SCR).	PTS system, sucrose- specific IIBC component (EIIBC-SCR) (Sucrose- permease IIBC component) (Phosphotransferase enzyme II, BC component) (EC 2.7.1.69) (EII-SCR).	PTS system, sucrose- specific IIBC component (EIIBC-SCR) (Sucrose- permease IIBC component) (Phosphotransferase enzyme II, BC component) (EC 2.7.1.69) (EII-SCR).
VIBRIO ALGINOLYTICUS. 01-AUG-1991 BACTERIA; PROTEOBACTERIA; GAMMAPROTEOBACTERI A; VIBRIONALES; VIBRIONACEAE; VIBRIO.	STAPHYLOCOCCUS XYLOSUS. BACTERIA; FIRMICUTES; BACILLALES; STAPHYLOCOCCUS.	SALMONELLA TYPHIMURIUM. BACTERIA; PROTEOBACTERIA; GAMMAPROTEOBACTERI A; ENTEROBACTERIALES; ENTEROBACTERIACEAE; SALMONELLA.
01-AUG-1991	01-OCT-1996	01-AUG-1988

Search term = sucrose symporter

Sucrose transpoter coding sequences available in EMBL or SwissProt databases prior to August 07, 1997



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Description:symporter*]) | [libs={embl swissprot}-Description:sucrose symporter*]) & [libs-DateCreated#19000101:19970807]) " found 6 entries Query "((([libs={embl swissprot}-Description:sucrose*] & [libs={embl swissprot}-

EXIBL Swiss-Prot	IE	<u>Acc Number</u>	Molecule	Description	<u> Urganism</u>
EMBL:HSZZ29997	HSZZ29997	AA324944	RNA	HSZZ29997 AA324944 RNA EST27854 Cerebellum II Homo sapiens cDNA 5' end similar to similar to sucrose/proton symporter.	Homo sapiens (human) Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
FMBL:ATSUC2S	ATSUC2S X75382	X75382	RNA	RNA A.thaliana SUC2 mRNA for sucrose- proton symporter	Arabidopsis thaliana (thale cress) Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids II; Brassicales; Brassicaceae; Arabidopsis.
F EMBL:ATSUCIS	ATSUCIS X75365	:	RNA	RNA A.thaliana SUCI	Arabidopsis thaliana (thale cress)

B.vulgaris mRNA for Beta vulgaris subsp. vulgaris sucrose/proton- Eukaryota; Viridiplantae; Streptophyta:
